

(19) United States

(12) Patent Application Publication (10) Pub. No.: US 2021/0302433 A1 Chen et al.

Sep. 30, 2021 (43) **Pub. Date:**

(54) METHODS FOR CHARACTERIZING LOW-ABUNDANCE HOST CELL PROTEINS

(71) Applicant: Regeneron Pharmaceuticals, Inc., Tarrytown, NY (US)

(72) Inventors: I-Hsuan Chen, Redwood City, CA (US); Hui Xiao, Scarsdale, NY (US); Ning Li, New Canaan, CT (US)

(21) Appl. No.: 17/217,518

(22) Filed: Mar. 30, 2021

Related U.S. Application Data

(60) Provisional application No. 63/001,690, filed on Mar. 30, 2020, provisional application No. 63/031,336, filed on May 28, 2020.

Publication Classification

(51) Int. Cl. G01N 33/68

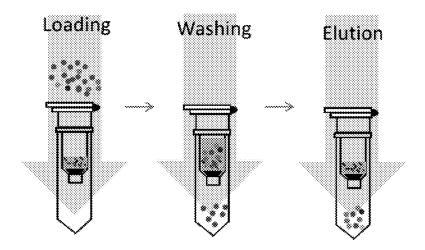
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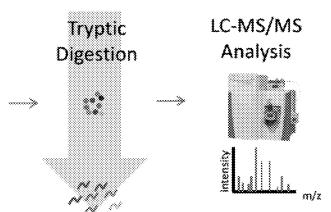
U.S. Cl.

CPC G01N 33/6848 (2013.01); G01N 2560/00 (2013.01); G01N 2458/15 (2013.01)

(57)**ABSTRACT**

The present application provides methods and systems to identify host cell protein (HCP) impurities in a sample containing high-abundance proteins. The HCP impurities can be enriched using interacting peptide ligands which have been attached to solid support. The HCP impurities can be eluted from the solid support using solution containing phase transfer surfactants. The isolated HCP impurities can be digested to generate components of the isolated HCP impurities which can subsequently be identified using a mass spectrometer.





- Proteominer beads
- Therapeutic mAb

